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CXMS analysis of model proteins

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An abbreviated version of this protocol was published in eLIFE in Mar 2016

Trifunctional cross-linker for mapping protein-protein interaction networks and comparing protein conformational states

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1. Dong, M. (2021). CXMS analysis of model proteins. Bio-protocol Preprint. bio-protocol.org/prep1358.
2. Tan, D., Li, Q., Zhang, M., Liu, C., Ma, C., Zhang, P., Ding, Y., Fan, S., Tao, L., Yang, B., Li, X., Ma, S., Liu, J., Feng, B., Liu, X., Wang, H., He, S., Gao, N., Ye, K., Dong, M. and Lei, X.(2016). Trifunctional cross-linker for mapping protein-protein interaction networks and comparing protein conformational states. eLIFE. DOI: 10.7554/eLife.12509

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